

SEQUENCE LISTING

<110> Klein, Elliott S.
Chandraratna Roshantha A.

<120> Methods of Detecting Dissociated Nuclear
Hormone Receptor Ligands

<130> P-AR 4528

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<170> FastSEQ for Windows Version 4.0

<210> 1

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<212> PRT

<213> Homo sapiens

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Asn	Gly	Tyr	Pro	Val	Pro	Pro	Tyr	Ala	Phe	Phe	Phe	Pro	Pro	Met	Leu
			20					25						30	
Gly	Gly	Leu	Ser	Pro	Pro	Gly	Ala	Leu	Thr	Thr	Leu	Gln	His	Gln	Leu
			35				40					45			
Pro	Val	Ser	Gly	Tyr	Ser	Thr	Pro	Ser	Pro	Ala	Thr	Ile	Glu	Thr	Gln
			50			55					60				
Ser	Ser	Ser	Ser	Glu	Glu	Ile	Val	Pro	Ser	Pro	Pro	Ser	Pro	Pro	Pro
65					70					75					80
Leu	Pro	Arg	Ile	Tyr	Lys	Pro	Cys	Phe	Val	Cys	Gln	Asp	Lys	Ser	Ser
				85				90						95	
Gly	Tyr	His	Tyr	Gly	Val	Ser	Ala	Cys	Glu	Gly	Cys	Lys	Gly	Phe	Phe
			100					105						110	
Arg	Arg	Ser	Ile	Gln	Lys	Asn	Met	Val	Tyr	Thr	Cys	His	Arg	Asp	Lys
			115				120					125			
Asn	Cys	Ile	Ile	Asn	Lys	Val	Thr	Arg	Asn	Arg	Cys	Gln	Tyr	Cys	Arg
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Leu	Gln	Lys	Cys	Phe	Glu	Val	Gly	Met	Ser	Lys	Glu	Ser	Val	Arg	Asn
145					150					155					160
Asp	Arg	Asn	Lys	Lys	Lys	Lys	Glu	Val	Pro	Lys	Pro	Glu	Cys	Ser	Glu
				165					170					175	
Ser	Tyr	Thr	Leu	Thr	Pro	Glu	Val	Gly	Glu	Leu	Ile	Glu	Lys	Val	Arg
			180					185					190		
Lys	Ala	His	Gln	Glu	Thr	Phe	Pro	Ala	Leu	Cys	Gln	Leu	Gly	Lys	Tyr
			195				200					205			
Thr	Thr	Asn	Asn	Ser	Ser	Glu	Gln	Arg	Val	Ser	Leu	Asp	Ile	Asp	Leu
			210			215					220				
Trp	Asp	Lys	Phe	Ser	Glu	Leu	Ser	Thr	Lys	Cys	Ile	Ile	Lys	Thr	Val
225					230					235					240
Asp	Phe	Ala	Lys	Gln	Leu	Pro	Gly	Phe	Thr	Thr	Leu	Thr	Ile	Ala	Asp

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                245                250                255
Gln Ile Thr Leu Leu Lys Ala Ala Cys Leu Asp Ile Leu Ile Leu Arg
                260                265                270
Ile Cys Thr Arg Tyr Thr Pro Glu Gln Asp Thr Met Thr Phe Ser Asp
                275                280                285
Gly Leu Thr Leu Asn Arg Thr Gln Met His Asn Ala Gly Phe Gly Pro
                290                295                300
Leu Thr Asp Leu Val Phe Ala Phe Ala Asn Gln Leu Leu Pro Leu Glu
305                310                315                320
Met Asp Asp Ala Glu Thr Gly Leu Leu Ser Ala Ile Cys Leu Ile Cys
                325                330                335
Gly Asp Arg Gln Asp Leu Glu Gln Pro Asp Arg Val Asp Met Leu Gln
                340                345                350
Glu Pro Leu Leu Glu Ala Leu Lys Val Tyr Val Arg Lys Arg Arg Pro
                355                360                365
Ser Arg Pro His Met Phe Pro Lys Met Leu Met Lys Ile Thr Asp Leu
                370                375                380
Arg Ser Ile Ser Ala Lys Gly Ala Glu Arg Val Ile Thr Leu Lys Met
385                390                395                400
Glu Ile Pro Gly Ser Met Pro Pro Leu Ile Gln Glu Met Leu Glu Asn
                405                410                415
Ser Glu Gly Leu Asp Thr Leu Ser Gly Gln Pro Gly Gly Gly Gly Arg
                420                425                430
Asp Gly Gly Gly Leu Ala Pro Pro Pro Gly Ser Cys Ser Pro Ser Leu
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Ser Pro Ser Ser Asn Arg Ser Ser Pro Ala Thr His Ser Pro
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<213> Homo sapiens

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Leu Lys Ala Cys Phe Ser Gly Leu Thr Gln Thr Glu Trp Gln His Arg
                35                40                45
His Thr Ala Gln Ser Ile Glu Thr Gln Ser Thr Ser Ser Glu Glu Leu
                50                55                60
Val Pro Ser Pro Pro Ser Pro Leu Pro Pro Pro Arg Val Tyr Lys Pro
65                70                75                80
Cys Phe Val Cys Gln Asp Lys Ser Ser Gly Tyr His Tyr Gly Val Ser
                85                90                95
Ala Cys Glu Gly Cys Lys Gly Phe Phe Arg Arg Ser Ile Gln Lys Asn
                100                105                110
Met Ile Tyr Thr Cys His Arg Asp Lys Asn Cys Val Ile Asn Lys Val
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Thr Arg Asn Arg Cys Gln Tyr Cys Arg Leu Gln Lys Cys Phe Glu Val
                130                135                140

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Gly Met Ser Lys Glu Ser Val Arg Asn Asp Arg Asn Lys Lys Lys Lys
 145 150 155 160
 Glu Thr Ser Lys Gln Glu Cys Thr Glu Ser Tyr Glu Met Thr Ala Glu
 165 170 175
 Leu Asp Asp Leu Thr Glu Lys Ile Arg Lys Ala His Gln Glu Thr Phe
 180 185 190
 Pro Ser Leu Cys Gln Leu Ala Lys Tyr Thr Thr Asn Ser Ser Ala Asp
 195 200 205
 His Arg Val Arg Leu Asp Leu Gly Leu Trp Asp Lys Phe Ser Glu Leu
 210 215 220
 Ala Thr Lys Cys Ile Ile Lys Ile Val Glu Phe Ala Lys Arg Leu Pro
 225 230 235 240
 Gly Phe Thr Gly Leu Thr Ile Ala Asp Gln Ile Thr Leu Leu Lys Ala
 245 250 255
 Ala Cys Leu Asp Ile Leu Ile Leu Arg Ile Cys Thr Arg Tyr Thr Pro
 260 265 270
 Glu Gln Asp Thr Met Thr Phe Ser Asp Gly Leu Thr Leu Asn Arg Thr
 275 280 285
 Gln Met His Asn Ala Gly Phe Gly Pro Leu Thr Asp Leu Val Phe Thr
 290 295 300
 Phe Ala Asn Gln Leu Leu Pro Leu Glu Met Asp Asp Thr Glu Thr Gly
 305 310 315 320
 Leu Leu Ser Ala Ile Cys Leu Ile Cys Gly Asp Arg Gln Asp Leu Glu
 325 330 335
 Glu Pro Thr Lys Val Asp Lys Leu Gln Glu Pro Leu Leu Glu Ala Leu
 340 345 350
 Lys Ile Tyr Ile Arg Lys Arg Arg Pro Ser Lys Pro His Met Phe Pro
 355 360 365
 Lys Ile Leu Met Lys Ile Thr Asp Leu Arg Ser Ile Ser Ala Lys Gly
 370 375 380
 Ala Glu Arg Val Ile Thr Leu Lys Met Glu Ile Pro Gly Ser Met Pro
 385 390 395 400
 Pro Leu Ile Gln Glu Met Met Glu Asn Ser Glu Gly His Glu Pro Leu
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 Thr Pro Ser Ser Ser Gly Asn Thr Ala Glu His Ser Pro Ser Ile Ser
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 Pro Ser Ser Val Glu Asn Ser Gly Val Ser Gln Ser Pro Leu Val Gln
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 Gly Ser Gly Tyr Pro Gly Ala Gly Phe Pro Phe Ala Phe Pro Gly Ala
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 Leu Arg Gly Ser Pro Pro Phe Glu Met Leu Ser Pro Ser Phe Arg Gly
 35 40 45
 Leu Gly Gln Pro Asp Leu Pro Lys Glu Met Ala Ser Leu Ser Val Glu

Gly Leu Lys Ser Pro Ala
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<210> 4
<211> 9
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<213> Artificial Sequence

<220>
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<222> (1)...(9)
<223> Xaa = Any Amino Acid

<221> VARIANT
<222> (1)...(9)
<223> Xaa = Any Amino Acid

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<210> 6
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His His His His His His
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15

<210> 17

<211> 27

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<220>

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<400> 17

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27

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<212> PRT

<213> Homo sapiens

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Lys	Arg	Thr	Val	Arg	Xaa	Asp	Leu	Thr	Tyr	Thr	Cys	Arg	Asp	Asn	Lys
			20					25					30		
Asp	Cys	Leu	Ile	Asp	Lys	Arg	Gln	Arg	Asn	Arg	Cys	Gln	Tyr	Cys	Arg
		35					40					45			
Tyr	Gln	Lys	Cys	Leu	Ala	Met	Gly	Met							
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<212> PRT

<213> T. cystophoro

<400> 19

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Lys	Arg	Ser	Val	Arg	Asn	Asn	Arg	Lys	Tyr	Ser	Cys	Leu	Gly	Lys	Arg
			20					25					30		
His	Cys	Asp	Thr	Asp	Lys	Lys	Ser	Arg	Asn	Arg	Cys	Gln	Tyr	Cys	Arg
		35					40					45			
Phe	Gln	Lys	Cys	Val	Gln	Val	Gly	Met							
		50				55									

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<213> Homo sapiens

<400> 20

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			20					25					30		
Glu	Gly	Lys	Cys	Val	Ile	Asp	Lys	Val	Thr	Arg	Asn	Gln	Cys	Gln	Glu
		35					40				45				
Cys	Arg	Phe	Lys	Lys	Cys	Ile	Tyr	Val	Gly	Met	Ala	Thr	Asp	Leu	Val
	50					55				60					
Leu	Asp	Gln	Ser	Lys	Arg	Leu	Ala	Lys	Arg	Lys	Leu	Ile	Glu	Glu	Asn
65					70				75					80	
Arg	Glu	Lys													

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 Gln Cys Thr Ile Asp Lys Asn Arg Arg Lys Ser Cys Gln Ala Cys Arg
 35 40 45
 Leu Arg Lys Cys Tyr Glu Val Gly Met Met Lys Gly Gly
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 <212> PRT
 <213> Rattus sp.

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 35 40 45
 Tyr Arg Lys Cys Leu Gln Ala Gly Met
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<210> 23
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 1 5 10 15
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<213> Homo sapiens

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Phe Ala Lys Lys Leu Pro Met Phe Ser Glu Leu Pro Cys Glu Asp Gln
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Ile Ile Leu Leu
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<213> Homo sapiens

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Ile Thr Leu Leu
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Tyr Ala Lys Ser Ile Pro Gly Phe Val Asn Leu Asp Leu Asn Asp Gln
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Val Thr Leu Leu
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<213> Homo sapiens

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Ile Val Leu Leu
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Met Thr Leu Leu
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<210> 44
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1 5

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Met Val Ser Glu Val Ile
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